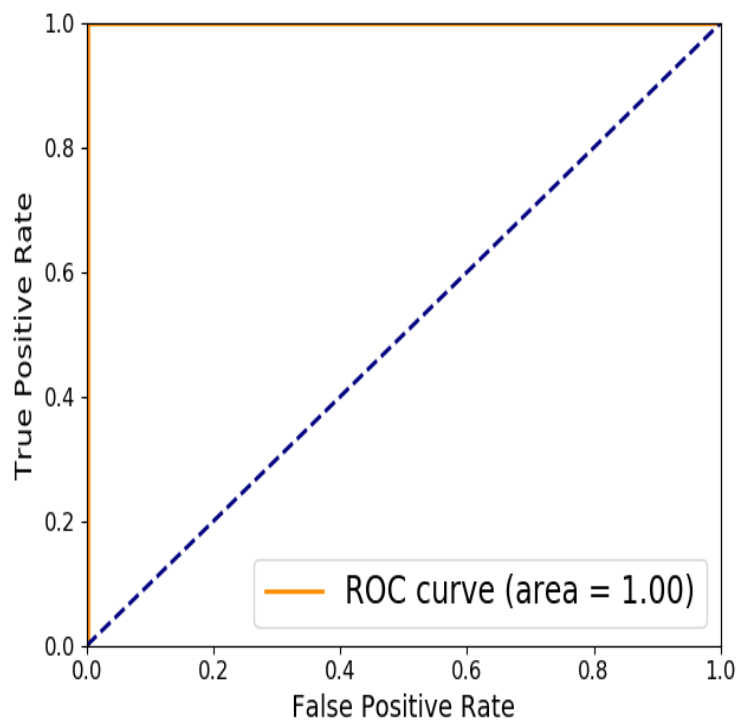
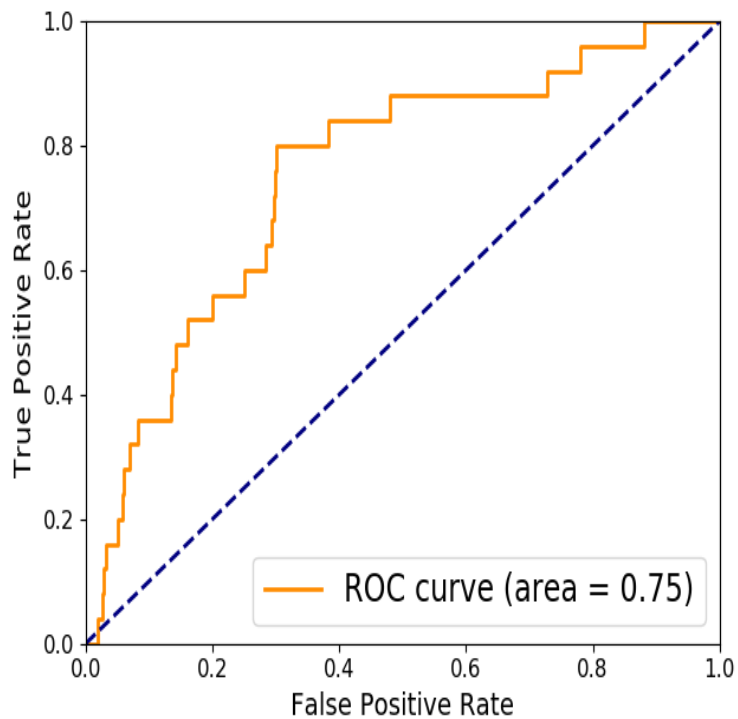


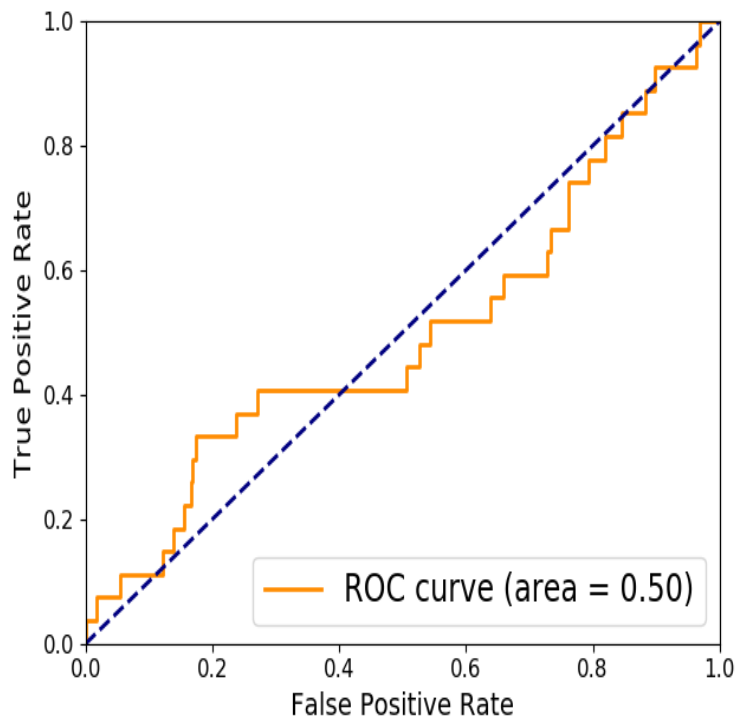
Diffusion ROC from fGOnly_tOverlap STRING10_combined.png



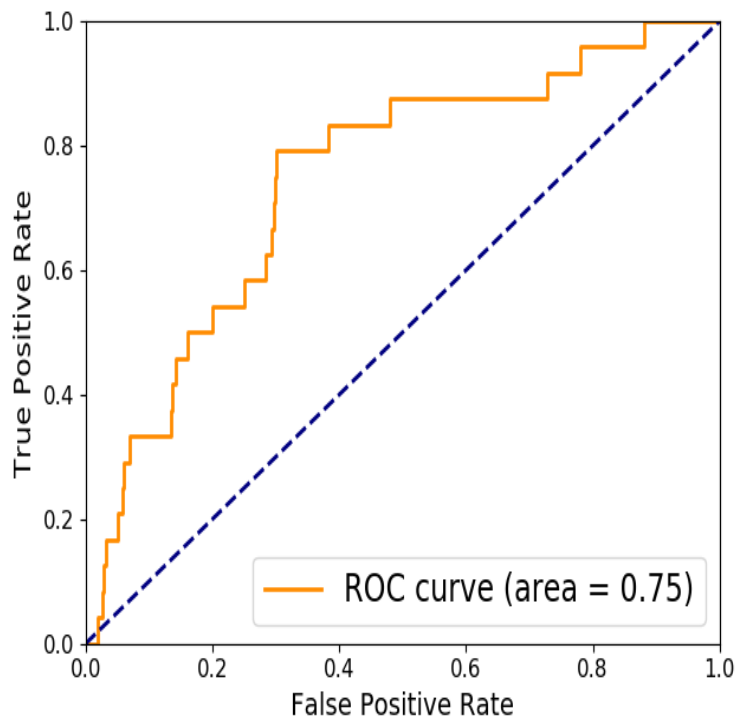
Diffusion ROC from fTestOnly_tGS STRING10_combined.png



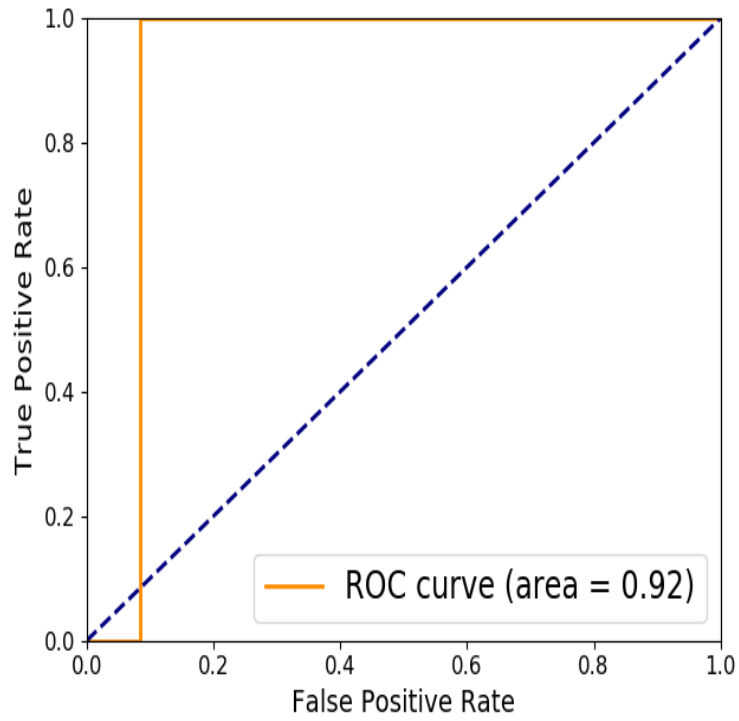
Diffusion ROC from fGOnly_tTestOnly STRING10_combined.png



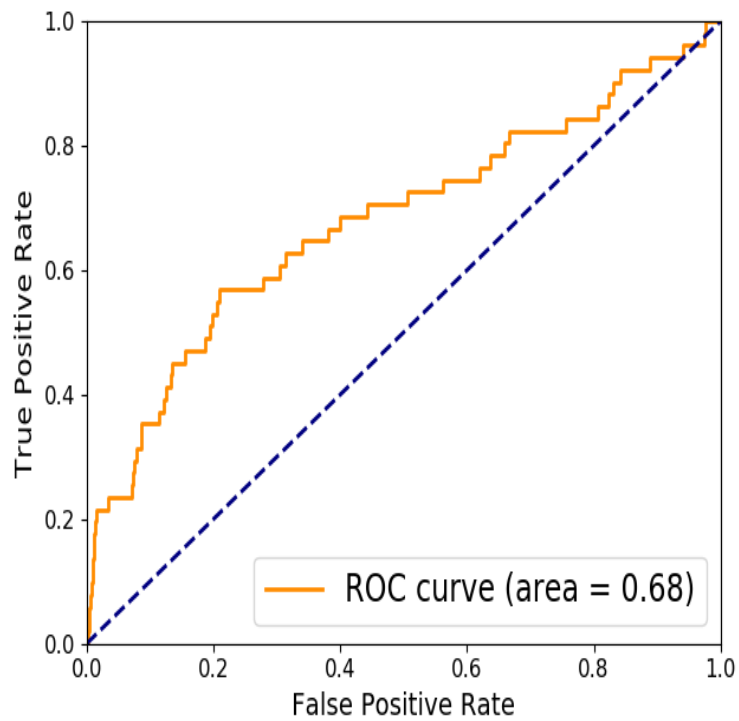
Diffusion ROC from fTestOnly_tGOnly STRING10_combined.png



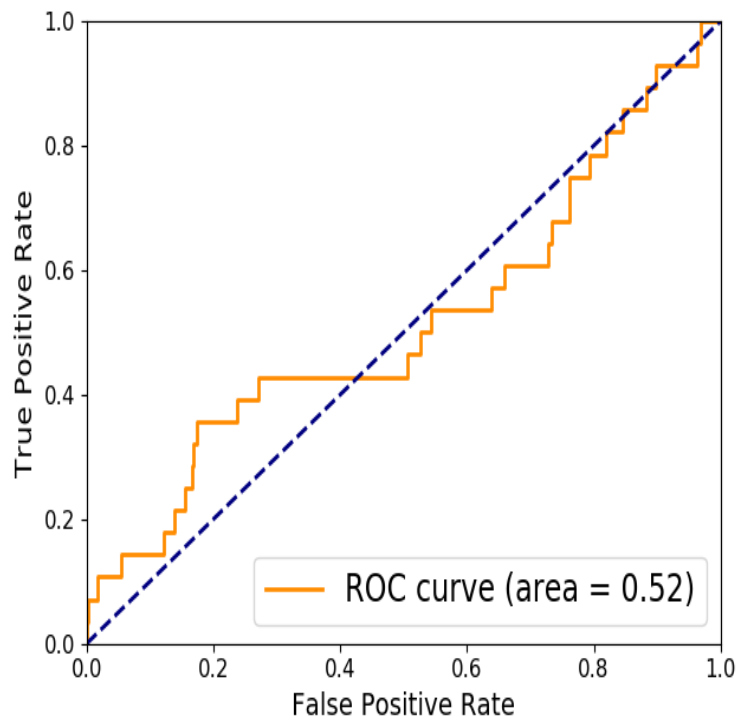
Diffusion ROC from fTestOnly_tOverlap STRING10_combined.png



Diffusion ROC from fOverlap_tOnlies STRING10_combined.png



Diffusion ROC from fGOnly_tTest STRING10_combined.png



Diff_FROM	Diff_TO	AUC	Zt_wdegre	Zt	Zf_wdegre	Zf
		e		e		
GS_only	Test_only	0.499	1.633	-0.017925	1.511	1.100052
			866278734		58115674	
			298		45 (inva	
			(invalid)		lid)	
Test_only	GS_only	0.746	3.509	4.634	2.6540086	2.866
					774943665	
					(invalid)	
GS_only	Overlap	0.998	5.3053892	1.5961048	1.7393261	1.874959
			22082631	180100876	976893581	72232287
			(invalid)	(invalid)	(invalid)	93 (inva
						lid)
GS_only	Test	0.517	1.537	0.422	1.488	1.688
Test_only	Overlap	0.916	2.2880637	1.4488158	1.208	0.864904
			96910328	634878163		93543474
			(invalid)	(invalid)		26 (inva
						lid)
Test_only	GS	0.753	4.101	4.204	2.756	2.256599
						18573086
						3 (inval
						id)
Overlap	Onlies	0.680	5.262	4.314	6.1793518	2.148
						52061758
						(invalid)

**	#Mapped	#Total	Not mapped genes
GS	24	24	N/A
Test_only	27	27	N/A
Overlap	1	1	APOE

(end of file)